# 1056/236

# SEQUENCE LISTING

## SEQ ID NO: 1

MVLLYSQASWDKRSKADALVLPFWMKNSKAQEAAVVDEDYKLVYQNALSNFSGKKG ETAFLFGNDHTKEQKIVLLGLGKSEEVSGTTVLEAYAQATTVLRKAKCKTVNILLPTISQ LRFSVEEFLTNLAAGVLSLNYNYPTYHKVDTSLPFLEKVTVMGIVSKVGDKIFRKEESLF EGVYLTRDLVNTNADEVTPEKLAAVAKDLAGEFASLDVKILDRKAILKEKMGLLAAVA KGAAVEPRFIVLDYQGKPKSKDRTVLIGKGVTFDSGGLDLKPGKAMITMKEDMAGAAT VLGIFSALASLELPINVTGIIPATENAIGSAAYKMGDVYVGMTGLSVEIGSTDAEGRLILA DAISYALKYCNPTRIIDFATLTGAMVVSLGESVAGFFANNDVLARDLAEASSETGEALW RMPLVEKYDQALHSDIADMKNIGSNRAGSITAALFLQRFLEDNPVAWAHLDIAGTAYHE KEELPYPKYATGFGVRCLIHYMEKFLSK

#### SEQ ID NO: 2

MTASGGAGGLGSTQTVDVARAQAAAATQDAQEVIGSQEASEASMLKGCEDLINPAAAT RIKKKGEKFESLEARRKPTADKAEKKSESTEEKGDTPLEDRFTEDLSEVSGEDFRGLKNS FDDDSSPDEILDALTSKFSDPTIKDLALDYLIQTAPSDGKLKSTLIQAKHQLMSQNPQAIV GGRNVLLASETEASRANTSPSSLRSLYFQVTSSPSNCANLHQMLASYLPSEKTAVMEFLV NGMVADLKSEGPSIPPAKLQVYMTELSNLQALHSVNSFFDRNIGNLENSLKHEGHAPIPS LTTGNLTKTFLQLVEDKFPSSSKAQKALNELVGPDTGPQTEVLNLFFRALNGCSPRIFSGA EKKQQLASVITNTLDAINADNEDYPKPGDFPRSSFSSTPPHAPVPQSEIPTSPTSTQPPSP

#### SEQ ID NO: 3

MCIKRKKTWIAFLAVVCSFCLTGCLKEGGDSNSEKFIVGTNATYPPFEFVDKRGEVVGFD IDLAREISNKLGKTLDVREFSFDALILNLKQHRIDAVITGMSITPSRLKEILMIPYYGEEIKH LVLVFKGENKHPLPLTQYRSVAVQTGTYQEAYLQSLSEVHIRSFDSTLEVLMEVMHGKS PVAVLEPSIAQVVLKDFPALSTATIDLPEDQWVLGYGIGVASDRPALALKIEAAVQEIRKE GVLAELEQKWGLNN

#### SEQ ID NO: 4

MSEKRKSNKIIGIDLGTTNSCVSVMEGGQPKVIASSEGTRTTPSIVAFKGGETLVGIPAKR QAVTNPEKTLASTKRFIGRKFSEVESEIKTVPYKVAPNSKGDAVFDVEQKLYTPEEIGAQI LMKMKETAEAYLGETVTEAVITVPAYFNDSQRASTKDAGRIAGLDVKRIIPEPTAAALA YGIDKEGDKKIAVFDLGGGTFDISILEIGDGVFEVLSTNGDTHLGGDDFDGVIINWMLDEF KKQEGIDLSKDNMALQRLKDAAEKAKIELSGVSSTEINQPFITIDANGPKHLALTLTRAQF EHLASSLIERTKQPCAQALKDAKLSASDIDDVLLVGGMSRMPAVQAVVKEIFGKEPNKG VNPDEVVAIGAAIQGGVLGGEVKDVLLLDVIPLSLGIETLGGVMTPLVERNTTIPTQKKQI FSTAADNQPAVTIVVLQGERPMAKDNKEIGRFDLTDIPPAPRGHPQIEVTFDIDANGILHV SAKDAASGREQKIRIEASSGLKEDEIQQMIRDAELHKEEDKQRKEASDVKNEADGMIFRA EKAVKDYHDKIPAELVKEIEEHIEKVRQAIKEDASTTAIKAASDELSTHMQKIGEAMQAQ SASAAASSAANAQGGPNINSEDLKKHSFSTRPPAGGSASSTDNIEDADVEIVDKPE

## SEQ ID NO: 5

MHDALQSILAIQELDIKMIRLMRVKKEHQNELAKIQALKTDIRRKVEEKEQEMEKLKDQI KGGEKRIQEISDQINKLENQQAAVKKMDEFNALTQEMTAANKERRTLEHQLSDLMDKQ AGSEDLLISLKESLSSTENSSSAIEEEIRENIRKINEEGRSLLSQRTQLKETTDPELFSIYERL LNNKKDRVVVPIENRVCSGCHIALTPQHENLVRKQDHLVFCEHCSRILYWQELQSPSAE GATTKRRRRRTAV

SEQ ID NO: 6

MKKFLLLSLMSLSSLPTFAANSTGTIGIVNLRRCLEESALGKKESAEFEKMKNQFSNSMG KMEEELSSIYSKLQDDDYMEGLSETAAAELRKKFEDLSAEYNTAQGQYYQILNQSNLKR MQKIMEEVKKASETVRIQEGLSVLLNEDIVLSIDSSADKTDAVIKVLDDSFQNN

# SEQ ID NO: 7

MTTESLETLVEQLSGLTVLELSQLKKLLEEKWDVTAAAPVVAVAGAAAAGDAPASAEP TEFAVILEDVPSDKKIGVLKVVREVTGLALKEAKEMTEGLPKTVKEKTSKSDAEDTVKK LQEAGAKAVAKGL

## SEQ ID NO: 8

MKKTALLAALCSVVSLSSCCRIVDCCFEDPCAPIQCSPCESKKKDVDGGCNSCNGYVPA CKPCGGDTHQDAKHGPQARGIPVDGKCRQ

## SEQ ID NO: 9

MPKIDTCDSCVSNTELLAIRTRVTQSYNEAQTILSSIPDGIFLLSESGEILICNPQARAILGIP EDIQLVTRMFHDFFPDTFFGFSVQEALEKEVPPKTIRLTLSQELSQKEVEVFVRKNISHDFL FLLIRDRSDYRQLEQAIEKYRSISELGKIAATLAHEIRNPLTSISGFATLLKEELSSERHQRM LNVIEGTRSLNSLVSSMLEYTKIQPLNLRSIDLQDFFSSLIPELSLTFPSCTFRRTILSPIQRSI DPDRLRCVIWNLVKNAVEASDEEIFLELHEKGFSVINTGTLPPNIQEKLFIPFFTTKPQGNG LGLAEAHKIMRLHGGDLVVSTQDNRTTFTILWTPA

#### SEQ ID NO: 10

MKVILRALCLFLVLPCGCYARVPSFEPFRGAIAPNRYTPKHSPELYFEMGDKYFQAKKFK QALLCFGMITHHFPEHALHPKAQFLVGLCYLEMGHPDLADKALTQYQELADTEYSEQLF AIKYSIAQSFANGKRKNIVPLEGFPKLLKADTDALRIFEEIVTASSDADLKASALYAKGAL LFDRKEYSEAIKTLKKVSLQFPSHSLSPESFTLIAKIHCLQALQEPYNEQYLQDARMNAAA LRKQHPNHPSNTEVENYIHHMCEAYASCLYSTGRFYEKKRKASSAKIYYSIALENFPDTS YVAKCNKRLERLSKQMS

#### SEQ ID NO: 11

MFDVVISDIEAREILDSRGYPTLCVKVITNTGTFGEACVPSGASTGIKEALELRDKDPKRY QGKGVLQAISNVEKVLMPALQGFSVFDQITADAIMIDADGTPNKEKLGANAILGVSLAL AKAAANTLQRPLYRYLGGSFSHVLPCPMMNLINGGMHATNGLQFQEFMIRPISAPSLTE AVRMGAEVFNALKKILQNRQLATGVGDEGGFAPNLASNAEALDLLLTAIETAGFTPRED ISLALDCAASSFYNTQDKTYDGKSYADQVGILAELCEHYPIDSIEDGLAEEDFEGWKLLS ETLGDRVQLVGDDLFVTNSALIAEGIAQGLANAVLIKPNQIGTLTETAEAIRLATIQGYAT ILSHRSGETEDTTIADLAVAFNTGQIKTGSLSRSERIAKYNRLMAIEEEMGPEALFQDSNFF SKA

#### SEQ ID NO: 12

MMKRLLCVLLSTSVFSSPMLGYSASKKDSKADICLAVSSGDQEVSQEDLLKEVSRGFSR VAAKATPGVVYIENFPKTGNQAIASPGNKRGFQENPFDYFNDEFFNRFFGLPSHREQQRP QQRDAVRGTGFIVSEDGYVVTNHHVVEDAGKIHVTLHDGQKYTAKIVGLDPKTDLAVI KIQAEKLPFLTFGNSDQLQIGDWAIAIGNPFGLQATVTVGVISAKGRNQLHIVDFEDFIQT DAAINPGNSGGPLLNINGQVIGVNTAIVSGSGGYIGIGFAIPSLMAKRVIDQLISDGQVTRG FLGVTLQPIDSELATCYKLEKVYGALVTDVVKGSPAEKAGLRQEDVIVAYNGKEVESLS ALRNAISLMMPGTRVVLKIVREGKTIEIPVTVTQIPTEDGVSALQKMGVRVQNITPEICKK LGLAADTRGILVVAVEAGSPAASAGVAPGQLILAVNRQRVASVEELNQVLKNSKGENVL LMVSQGDVVRFIVLKSDE

**SEQ ID NO: 13** 

MKKINKIVLAVGGTGGHIIPALAARETFIHEDIEVLLLGKGLAHFLGDDSEVAYCDIPSGS PFSLRVNRMFSGAKQLYKGYVAALQKIRDFTPDLAIGFGSYHSLPAMLASIRSRIPLFLHE QNIVPGKVNKLFSRFAKGVGMSFAAAGEHFHCRAEEVFLPIRKLSEQIVFPGASPVICVV GGSQGAKILNDVVPKALARIRESYSNLYVHHIVGPKGDLQAVSQVYQDAGINHTVTAFD HNMLGVLQASDLVISRSGATMLNELLWVQVPAILIPYPGAYGHQEVNAKFFTHTVGGGT MILQKYLTEESLSKQVLLALDPATSENRRKAMLSAQQKKSFKSLYQFICESL

SEQ ID NO: 14

MGNSGFYLYNTQNCVFADNIKVGQMTEPLKDQQIILGTTSTPVAAKMTASDGISLTVSN NPSTNASITIGLDAEKAYQLILEKLGDQILGGIADTIVDSTVQDILDKITTDPSLGLLKAFN NFPITNKIQCNGLFTPRNIETLLGGTEIGKFTVTPKSSGSMFLVSADIIASRMEGGVVLALV REGDSKPYAISYGYSSGVPNLCSLRTRIINTGLTPTTYSLRVGGLESGVVWVNALSNGNDI LGITNTSNVSFLEVIPQTNA

**SEQ ID NO: 15 (pmpA)(CT412)** MNRVIEIHAHYDQRQLSQSPNTNFLVHHPYLTLIPKFLLGALIVYAPYSFAEMELAISGHK QGKDRDTFTMISSCPEGTNYIINRKLILSDFSLLNKVSSGGAFRNLAGKISFLGKNSSASIH **FKHININGEGAGVESESSIEFTDLRKLVAEGSESTGGIFTAKEDISEKNNHHIAERNNITKGN** GGVIQLQGDMKGSVSFVDQRGAIIFTNNQAVTSSSMKHSGRGGAISGDFAGSRILFLNNQ QITFEGNSAVHGGAIYNKNGLVEFLGNAGPLAFKENTTIANGGAIYTSNFKANQQTSPILF SQNHANKKGGAIYAQYVNLEQNQDTIRFEKNTAKEGGGAITSSQCSITAHNTIIFSDNAA GDLGGGAILLEGKKPSLTLIAHSGNIAFSGNTMLHITKKASLDRHNSILIKEAPYKIQLAA NKNHSIHFFDPVMALSASSSPIQINAPEYETPFFSPKGMIVFSGANLLDDAREDVANRTSIF NQPVHLYNGTLSIENGAHLIVQSFKQTGGRISLSPGSSLALYTMNSFFHGNISSKEPLEING LSFGVDISPSNLQAEIRAGNAPLRLSGSPSIHDPEGLFYENRDTAASPYQMEILLTSDKIVDI SKFTTDSLVTNKQSGFQGAWHFSWQPNTINNTKQKILRASWLPTGEYVLESNRVGRAVP NSLWSTFLLLQTASHNLGDHLCNNRSLIPTSYFGVLIGGTGAEMSTHSSEEESFISRLGAT GTSIIRLTPSLTLSGGGSHMFGDSFVADLPEHITSEGIVQNVGLTHVWGPLTVNSTLCAAL DHNAMVRICSKKDHTYGKWDTFGMRGTLGASYTFLEYDQTMRVFSFANIEATNILQRA FTETGYNPRSFSKTKLLNIAIPIGIGYEFCLGNSSFALLGKGSIGYSRDIKRENPSTLAHLAM NDFAWTTNGCSVPTSAHTLANQLILRYKACSLYITAYTINREGKNLSNSLSCGGYVGF

**SEQ ID NO: 16 (pmpB)(CT413)** MKWLSATAVFAAVLPSVSGFCFPEPKELNFSRVGTSSSTTFTETVGEAGAEYTVSGNASF TKFTNIPTTDTTTPTNSNSSSSNGETASVSEDSDSTTTTPDPKGGGAFYNAHSGVLSFMTR SGTEGSLTLSEIKITGEGGAIFSQGELLFTDLTGLTIQNNLSQLSGGAIFGESTISLSGITKAT FSSNSAEVPAPVKKPTEPKAQTASETSGSSSSSGNDSVSSPSSSRAEPAAANLQSHFICATA TPAAQTDTETSTPSHKPGSGGAIYAKGDLTIADSQEVLFSINKATKDGGAIFAEKDVSFEN ITSLKVQTNGAEEKGGAIYAKGDLSIQSSKQSLFNSNYSKQGGGALYVEGDINFQDLEEIR IKYNKAGTFETKKITLPKAQASAGNADAWASSSPQSGSGATTVSNSGDSSSGSDSDTSET VPATAKGGGLYTDKNLSITNITGIIEIANNKATDVGGGAYVKGTLTCENSHRLQFLKNSS DKQGGGIYGEDNITLSNLTGKTLFQENTAKEEGGGLFIKGTDKALTMTGLDSFCLINNTS  ${\tt EKHGGGAFVTKE} {\tt EISQTYTSDVETIPGITPVHGETVITGNKSTGGNGGGVCTKRLALSNLQ}$ SISISGNSAAENGGGAHTCPDSFPTADTAEQPAAASAATSTPESAPVVSTALSTPSSSTVSS LTLLAASSQASPATSNKETQDPNADTDLLIDYVVDTTISKNTAKKGGGIYAKKAKMSRID QLNISENSATEIGGGICCKESLELDALVSLSVTENLVGKEGGGLHAKTVNISNLKSGFSFS NNKANSSSTGVATTASAPAAAAASLQAAAAAVPSSPATPTYSGVVGGAIYGEKVTFSQC SGTCQFSGNQAIDNNPSQSSLNVQGGAIYAKTSLSIGSSDAGTSYIFSGNSVSTGKSQTTGQIAGGAIYSPTVTLNCPATFSNNTASMATPKTSSEDGSSGNSIKDTIGGAIAGTAITLSGVS RFSGNTADLGAAIGTLANANTPSATSGSQNSITEKITLENGSFIFERNQANKRGAIYSPSVS IKGNNITFNQNTSTHDGSAIYFTKDATIESLGSVLFTGNNVTATQASSATSGQNTNTANY

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GAAIFGDPGTTQSSQTDAILTLLASSGNITFSNNSLQNNQGDTPASKFCSIAGYVKLSLQA AKGKTISFFDCVHTSTKKIGSTQNVYETLDINKEENSNPYTGTIVFSSELHENKSYIPQNAI LHNGTLVLKEKTELHVVSFEQKEGSKLIMKPGAVLSNQNIANGALVINGLTIDLSSMGTP QAGEIFSPPELRIVATTSSASGGSGVSSSIPTNPKRISAAAPSGSAATTPTMSENKVFLTGDL TLIDPNGNFYQNPMLGSDLDVPLIKLPTNTSDVQVYDLTLSGDLFPQKGYMGTWTLDSN PQTGKLQARWTFDTYRRWVYIPRDNHFYANSILGSQNSMIVVKQGLINNMLNNARFDDI AYNNFWVSGVGTFLAQQGTPLSEEFSYYSRGTSVAIDAKPRQDFILGAAFSKMVGKTKAIKKMHNYFHKGSEYSYQASVYGGKFLYFLLNKQHGWALPFLIQGVVSYGHIKHDTTTLY PSIHERNKGDWEDLGWLADLRISMDLKEPSKDSSKRITVYGELEYSSIRQKQFTEIDYDPRHFDDCAYRNLSLPVGCAVEGAIMNCNILMYNKLALAYMPSIYRNNPVCKYRVLSSNEAGQVICGVPTRTSARAEYSTQLYLGPFWTLYGNYTIDVGMYTLSQMTSCGARMIF

SEQ ID NO: 17 (pmpC)(CT414) MKFMSATAVFAAALSSVTEASSIQDQIKNTDCNVSKLGYSTSQAFTDMMLADNTEYRA ADSVSFYDFSTSSRLPRKHLSSSSEASPTTEGVSSSSSGETDEKTEEELDNGGIIYAREKLTI SESQDSLSNQSIELHDNSIFFGEGEVIFDHRVALKNGGAIYGEKEVVFENIKSLLVEVNIAV EKGGSVYAKERVSLENVTEATFSSNGGEQGGGGTYSEQDMLISDCNNVHFQGNAAGAT AVKOCLDEEMIVLLAECVDSLSEDTLDSTPETEQTESNGNQDGSSETEDTQVSESPESTPS PDDVLGKGGGIYTEKSLTITGITGTIDFVSNIATDSGAGVFTKENLSCTNTNSLQFLKNSA GQHGGGAYVTQTMSVTNTTSESITTPPLIGEVIFSENTAKGHGGGICTNKLSLSNLKTVTL TKNSAKESGGAIFTDLASIPITDTPESSTPSSSSPASTPEVVASAKINRFFASTAKPAAPSLTE AESDQTDQTETSDTNSDIDVSIENILNVAINQNTSAKKGGAIYGKKAKLSRINNLELSGNS SQDVGGGLCLTESVEFDAIGSLLSHYNSAAKEGGAIHSKTVTLSNLKSTFTFADNTVKAI VESTPEAPEEIPPVEGEESTATEDPNSNTEGSSANTNLEGSQGDTADTGTGDVNNESQDTS ··· DTGNAESEEQLQDSTQSNEENTLPNSNIDQSNENTDESSDSHTEEITDESVSSSSESGSSTP QDGGAASSGAPSGDQSISANACLAKSYAASTDSSPVSNSSGSEEPVTSSSDSDVTASSDNP DSSSSGDSAGDSEEPTEPEAGSTTETLTLIGGGAIYGETVKIENFSGQGIFSGNKAIDNTTE GSSSKSDVLGGAVYAKTLFNLDSGSSRRTVTFSGNTVSSQSTTGQVAGGAIYSPTVTIATP VVFSKNSATNNANNTTDTQRKDTFGGAIGATSAVSLSGGAHFLENVADLGSAIGLVPGT QNTETVKLESGSYYFEKNKALKRATIYAPVVSIKAYTATFNQNRSLEEGSAIYFTKEASIE  ${\tt SLGSVLFTGNLVTLTLSTTTEGTPATTSGDVTKYGAAIFGQIASSNGSQTDNLPLKLIASG}$ GNICFRNNEYRPTSSDTGTSTFCSIAGDVKLTMQAAKGKTISFFDAIRTSTKKTGTQATAY DTLDINKSEDSETVNSAFTGTILFSSELHENKSYIPQNVVLHSGSLVLKPNTELHVISFEQK EGSSLVMTPGSVLSNQTVADGALVINNMTIDLSSVEKNGIAEGNIFTPPELRIIDTTTGGSG GTPSTDSESNQNSDDTEEQNNNDASNQGESANGSSSPAVAAAHTSRTRNFAAAATATPT TTPTATTTTSNQVILGGEIKLIDPNGTFFQNPALRSDQQISLLVLPTDSSKMQAQKIVLTGD IAPQKGYTGTLTLDPDQLQNGTISVLWKFDSYRQWAYVPRDNHFYANSILGSQMLMVT VKQGLLNDKMNLARFEEVSYNNLWISGLGTMLSQVGTPTSEEFTYYSRGASVALDAKP AHDVIVGAAFSKMIGKTKSLKRENNYTHKGSEYSYQASVYGGKPFHFVINKKTEKSLPL LLQGVISYGYIKHDTVTHYPTIRERNKGEWEDLGWLTALRVSSVLRTPAQGDTKRITVY GELEYSSIRQKQFTETEYDPRYFDNCTYRNLAIPMGLAFEGELSGNDILMYNRFSVAYML SIYRNSPTCKYQVLSSGEGGEIICGVPTRNSARGEYSTQLYLGPLWTLYGSYTIEADAHTL **AHMMNCGARMTF** 

**SEQ ID NO: 18 (pmpD)(CT812)** 

MSSEKDIKSTCSKFSLSVVAAILASVSGLASCVDLHAGGQSVNELVYVGPQAVLLLDQIR DLFVGSKDSQAEGQYRLIVGDPSSFQEKDADTLPGKVEQSTLFSVTNPVVFQGVDQQDQ VSSQGLICSFTSSNLDSPRDGESFLGIAFVGDSSKAGITLTDVKASLSGAALYSTEDLIFEKI KGGLEFASCSSLEQGGACAAQSILIHDCQGLQVKHCTTAVNAEGSSANDHLGFGGGAFF VTGSLSGEKSLYMPAGDMVVANCDGAISFEGNSANFANGGAIAASGKVLFVANDKKTS FIENRALSGGAIAASSDIAFQNCAELVFKGNCAIGTEDKGSLGGGAISSLGTVLLQGNHGI

TCDKNESASQGGAIFGKNCQISDNEGPVVFRDSTACLGGGAIAAQEIVSIQNNQAGISFEG GKASFGGGIACGSFSSAGGASVLGTIDISKNLGAISFSRTLCTTSDLGQMEYQGGGALFGE NISLSENAGVLTFKDNIVKTFASNGKILGGGAILATGKVEITNNSEGISFTGNARAPOALPT QEEFPLFSKKEGRPLSSGYSGGGAILGREVAILHNAAVVFEQNRLQCSEEEATLLGCCGG GAVHGMDSTSIVGNSSVRFGNNYAMGQGVSGGALLSKTVQLAGNGSVDFSRNIASLGG GALQASEGNCELVDNGYVLFRDNRGRVYGGAISCLRGDVVISGNKGRVEFKDNIATRLY VEETVEKVEEVEPAPEQKDNNELSFLGRAEQSFITAANQALFASEDGDLSPESSISSEELA KRRECAGGAIFAKRVRIVDNQEAVVFSNNFSDIYGGAIFTGSLREEDKLDGQIPEVLISGN AGDVVFSGNSSKRDEHLPHTGGGAICTQNLTISQNTGNVLFYNNVACSGGAVRIEDHGN VLLEAFGGDIVFKGNSSFRAQGSDAIYFAGKESHITALNATEGHAIVFHDALVFENLEER KSAEVLLINSRENPGYTGSIRFLEAESKVPQCIHVQQGSLELLNGATLCSYGFKQDAGAK LVLAAGAKLKILDSGTPVQQGHAISKPEAEIESSSEPEGAHSLWIAKNAQTTVPMVDIHTI SVDLASFSSSQQEGTVEAPQVIVPGGSYVRSGELNLELVNTTGTGYENHALLKNEAKVPL MSFVASGDEASAEISNLSVSDLQIHVVTPEIEEDTYGHMGDWSEAKIQDGTLVISWNPTG YRLDPQKAGALVFNALWEEGAVLSALKNARFAHNLTAQRMEFDYSTNVWGFAFGGFR TLSAENLVAIDGYKGAYGGASAGVDIQLMEDFVLGVSGAAFLGKMDSQKFDAEVSRKG VVGSVYTGFLAGSWFFKGQYSLGETQNDMKTRYGVLGESSASWTSRGVLADALVEYRS LVGPVRPTEYALHENPYVEVSYASMKEPGETEQGREARSFEDASLTNITIPLGMKEELAEI KGQFSEVNSLGISYAWEAYRKVEGGAVQLLEAGFDWEGAPMDLPRQELRVALENNTE WSSYFSTVLGLTAFCGGFTSTDSKLGYEANTGLRLIF

SEQ ID NO: 19 (pmpE)(CT869) MKKAFFFFLIGNSLSGLAREVPSRIFLMPNSVPDPTKESLSNKISLTGDTHNLTNCYLDNL RYILAILQKTPNEGAAVTITDYLSFFDTQKEGIYFAKNLTPESGGAIGYASPNSPTVEIRDTI GPVIFENNTCCRLFTWRNPYAADKIREGGAIHAQNLYINHNHDVVGFMKNFSYVQGGAI STANTFVVSENQSCFLFMDNICIQTNTAGKGGAIYAGTSNSFESNNCDLFFINNACCAGG AIFSPICSLTGNRGNIVFYNNRCFKNVETASSEASDGGAIKVTTRLDVTGNRGRIFFSDNIT KNYGGAIYAPVVTLVDNGPTYFINNIANNKGGAIYIDGTSNSKISADRHAIIFNENIVTNV TNANGTSTSANPPRRNAITVASSSGEILLGAGSSQNLIFYDPIEVSNAGVSVSFNKEADQT GSVVFSGATVNSADFHQRNLQTKTPAPLTLSNGFLCIEDHAQLTVNRFTQTGGVVSLGN GAVLSCYKNGTGDSASNASITLKHIGLNLSSILKSGAEIPLLWVEPTNNSNNYTADTAATF SLSDVKLSLIDDYGNSPYESTDLTHALSSQPMLSISEASDNQLQSENIDFSGLNVPHYGWQ GLWTWGWAKTQDPEPASSATITDPQKANRFHRTLLLTWLPAGYVPSPKHRSPLIANTLW GNMLLATESLKNSAELTPSGHPFWGITGGGLGMMVYQDPRENHPGFHMRSSGYSAGMI AGQTHTFSLKFSQTYTKLNERYAKNNVSSKNYSCQGEMLFSLQEGFLLTKLVGLYSYGD HNCHHFYTQGENLTSQGTFRSQTMGGAVFFDLPMKPFGSTHILTAPFLGALGIYSSLSHF TEVGAYPRSFSTKTPLINVLVPIGVKGSFMNATHRPQAWTVELAYQPVLYRQEPGIAAQL LASKGIWFGSGSPSSRHAMSYKISQQTQPLSWLTLHFQYHGFYSSSTFCNYLNGEIALRF

SEQ ID NO: 20 (pmpF)(CT870)
MIKRTSLSFACLSFFYLSTISILQANETDTLQFRRFTFSDREIQFVLDPASLITAQNIVLSNLQ
SNGTGACTISGNTQTQIFSNSVNTTADSGGAFDMVTTSFTASDNANLLFCNNYCTHNKG
GGAIRSGGPIRFLNNQDVLFYNNISAGAKYVGTGDHNEKNRGGALYATTITLTGNRTLAF
INNMSGDCGGAISADTQISITDTVKGILFENNHTLNHIPYTQAENMARGGAICSRRDLCSIS
NNSGPIVFNYNQGGKGGAISATRCVIDNNKERIIFSNNSSLGWSQSSSASNGGAIQTTQGF
TLRNNKGSIYFDSNTATHAGGAINCGYIDIRDNGPVYFLNNSAAWGAAFNLSKPRSATN
YIHTGTGDIVFNNNVVFTLDGNLLGKRKLFHINNNEITPYTLSLGAKKDTRIYFYDLFQW
ERVKENTSNNPPSPTSRNTITVNPETEFSGAVVFSYNQMSSDIRTLMGKEHNYIKEAPTTL
KFGTLAIEDDAELEIFNIPFTQNPTSLLALGSGATLTVGKHGKLNITNLGVILPIILKEGKSP
PCIRVNPQDMTQNTGTGQTPSSTSSISTPMIIFNGRLSIVDENYESVYDSMDLSRGKAEQLI
LSIETTNDGQLDSNWQSSLNTSLLSPPHYGYQGLWTPNWITTTYTITLNNNSSAPTSATSI

AEQKKTSETFTPSNTTTASIPNIKASAGSGSGSASNSGEVTITKHTLVVNWAPVGYIVDPIR RGDLIANSLVHSGRNMTMGLRSLLPDNSWFALQGAATTLFTKQQKRLSYHGYSSASKG YTVSSQASGAHGHKFLLSFSQSSDKMKEKETNNRLSSRYYLSALCFEHPMFDRIALIGAA ACNYGTHNMRSFYGTKKSSKGKFHSTTLGASLRCELRDSMPLRSIMLTPFAQALFSRTEP ASIRESGDLARLFTLEQAHTAVVSPIGIKGAYSSDTWPTLSWEMELAYQPTLYWKRPLLN TLLIQNNGSWVTTNTPLAKHSFYGRGSHSLKFSHLKLFANYQAEVATSTVSHYINAGGA LVF

**SEQ ID NO: 21 (pmpG)(CT871)** MQTSFHKFFLSMILAYSCCSLSGGGYAAEIMIPQGIYDGETLTVSFPYTVIGDPSGTTVFSA GELTLKNLDNSIAALPLSCFGNLLGSFTVLGRGHSLTFENIRTSTNGAALSDSANSGLFTIE GFKELSFSNCNSLLAVLPAATTNNGSQTPTTTSTPSNGTIYSKTDLLLLNNEKFSFYSNLV SGDGGAIDAKSLTVQGISKLCVFQENTAQADGGACQVVTSFSAMANEAPIAFIANVAGV RGGGIAAVQDGQQGVSSSTSTEDPVVSFSRNTAVEFDGNVARVGGGIYSYGNVAFLNNG KTLFLNNVASPVYIAAEQPTNGQASNTSDNYGDGGAIFCKNGAQAAGSNNSGSVSFDGE GVVFFSSNVAAGKGGAIYAKKLSVANCGPVQFLGNIANDGGAIYLGESGELSLSADYGD IIFDGNLKRTAKENAADVNGVTVSSQAISMGSGGKITTLRAKAGHQILFNDPIEMANGNN QPAQSSEPLKINDGEGYTGDIVFANGNSTLYQNVTIEQGRIVLREKAKLSVNSLSQTGGSL YMEAGSTLDFVTPQPPQQPPAANQLITLSNLHLSLSSLLANNAVTNPPTNPPAQDSHPAII GSTTAGSVTISGPIFFEDLDDTAYDRYDWLGSNQKIDVLKLQLGTQPSANAPSDLTLGNE MPKYGYQGSWKLAWDPNTANNGPYTLKATWTKTGYNPGPERVASLVPNSLWGSILDIR SAHSAIQASVDGRSYCRGLWVSGVSNFFYHDRDALGQGYRYISGGYSLGANSYFGSSMF GLAFTEVFGRSKDYVVCRSNHHACIGSVYLSTKQALCGSYLFGDAFIRASYGFGNQHMK TSYTFAEESDVRWDNNCLVGEIGVGLPIVITPSKLYLNELRPFVQAEFSYADHESFTEEGD QARAFRSGHLMNLSVPVGVKFDRCSSTHPNKYSFMGAYICDAYRTISGTQTTLLSHQET WTTDAFHLARHGVIVRGSMYASLTSNIEVYGHGRYEYRDTSRGYGLSAGSKVRF

**SEQ ID NO: 22 (pmpH)(CT872)** MPFSLRSTSFCFLACLCSYSYGFASSPQVLTPNVTTPFKGDDVYLNGDCAFVNVYAGAE NGSIISANGDNLTITGQNHTLSFTDSQGPVLQNYAFISAGETLTLKDFSSLMFSKNVSCGE KGMISGKTVSISGAGEVIFWDNSVGYSPLSIVPASTPTPPAPAPAPAASSSLSPTVSDARKG SIFSVETSLEISGVKKGVMFDNNAGNFGTVFRGNSNNNAGSGGSGSATTPSFTVKNCKGK VSFTDNVASCGGGVVYKGTVLFKDNEGGIFFRGNTAYDDLGILAATSRDQNTETGGGG GVICSPDDSVKFEGNKGSIVFDYNFAKGRGGSILTKEFSLVADDSVVFSNNTAEKGGGAI YAPTIDISTNGGSILFERNRAAEGGAICVSEASSGSTGNLTLSASDGDIVFSGNMTSDRPGE RSAARILSDGTTVSLNASGLSKLIFYDPVVQNNSAAGASTPSPSSSSMPGAVTINQSGNGS VIFTAESLTPSEKLQVLNSTSNFPGALTVSGGELVVTEGATLTTGTITATSGRVTLGSGAS LSAVAGAANNNYTCTVSKLGIDLESFLTPNYKTAILGADGTVTVNSGSTLDLVMESEAE VYDNPLFVGSLTIPFVTLSSSSASNGVTKNSVTINDADAAHYGYQGSWSADWTKPPLAP DAKGMVPPNTNNTLYLTWRPASNYGEYRLDPQRKGELVPNSLWVAGSALRTFTNGLKE HYVSRDVGFVASLHALGDYILNYTQDDRDGFLARYGGFQATAASHYENGSIFGVAFGQ LYGQTKSRMYYSKDAGNMTMLSCFGRSYVDIKGTETVMYWETAYGYSVHRMHTQYF NDKTQKFDHSKCHWHNNNYYAFVGAEHNFLEYCIPTRQFARDYELTGFMRFEMAGGW SSSTRETGSLTRYFARGSGHNMSLPIGIVAHAVSHVRRSPPSKLTLNMGYRPDIWRVTPH CNMEIIANGVKTPIQGSPLARHAFFLEVHDTLYIHHFGRAYMNYSLDARRRQTAHFVSM **GLNRIF** 

SEQ ID NO: 23 (pmpI)
MRPDHMNFCCLCAAILSSTAVLFGQDPLGETALLTKNPNHVVCTFFEDCTMESLFPALC
AHASQDDPLYVLGNSYCWFVSKLHITDPKEALFKEKGDLSIQNFRFLSFTDCSSKESSPSII
HQKNGQLSLRNNGSMSFCRNHAEGSGGAISADAFSLQHNYLFTAFEENSSKGNGGAIQA

QTFSLSRNVSPISFARNRADLNGGAICCSNLICSGNVNPLFFTGNSATNGGAICCISDLNTS EKGSLSLACNQETLFASNSAKEKGGAIYAKHMVLRYNGPVSFINNSAKIGGAIAIQSGGS LSILAGEGSVLFQNNSQRTSDQGLVRNAIYLEKDAILSSLEARNGDILFFDPIVQESSSKES PLPSSLQASVTSPTPATASPLVIQTSANRSVIFSSERLSEEEKTPDNLTSQLQQPIELKSGRL VLKDRAVLSAPSLSQDPQALLIMEAGTSLKTSSDLKLATLSIPLHSLDTEKSVTIHAPNLSI QKIFLSNSGDENFYENVELLSKEQNNIPLLTLSKEQSHLHLPDGNLSSHFGYQGDWTFSW KDSDEGHSLIANWTPKNYVPHPERQSTLVANTLWNTYSDMQAVQSMINTIAHGGAYLF GTWGSAVSNLFYAHDSSGKPIDNWHHRSLGYLFGISTHSLDDHSFCLAAGQLLGKSSDSFITSTETTSYIATVQAQLATPLMKISAQACYNESIHELKTKYRSFSKEGFGSWHSVAVSGEV CASIPIVSNGSGLFSSFSIFSKLQGFSGTQDGFEESSGEIRSFSASSFRNISLPMGITFEKKSQ KTRNYYYFLGAYIQDLKRDVESGPVVLLKNAVSWDAPMANLDSRAYMFRLTNQRALH RLQTLLNVSYVLRGQSHSYSLDLGTTYRF

**SEQ ID NO: 24 (MOMP)(CT681)** 

MKKLLKSVLVFAALSSASSLQALPVGNPAEPSLMIDGILWEGFGGDPCDPCATWCDAIS MRVGYYGDFVFDRVLKTDVNKEFQMGAKPTTDTGNSAAPSTLTARENPAYGRHMQDA EMFTNAACMALNIWDRFDVFCTLGATSGYLKGNSASFNLVGLFGDNENQKTVKAESVP NMSFDQSVVELYTDTTFAWSVGARAALWECGCATLGASFQYAQSKPKVEELNVLCNA AEFTINKPKGYVGKEFPLDLTAGTDAATGTKDASIDYHEWQASLALSYRLNMFTPYIGV KWSRASFDADTIRIAQPKSATAIFDTTTLNPTIAGAGDVKTGAEGQLGDTMQIVSLQLNK MKSRKSCGIAVGTTIVDADKYAVTVETRLIDERAAHVNAQFRF

**SEQ ID NO: 25 CSFIGGITYL** 

SEQ ID NO: 26 SFIGGITYL

**SEQ ID NO: 27 SIIGGITYL** 

SEQ ID NO: 28 (Cap1)(CT529)

MASICGRLGSGTGNALKAFFTQPNNKMARVVNKTKGMDKTIKVAKSAAELTANILEQA GGAGSSAHITASQVSKGLGDARTVVALGNAFNGALPGTVQSAQSFFSHMKAASQKTQE GDEGLTADLCVSHKRRAAAAVCSIIGGITYLATFGAIRPILFVNKMLAKPFLSSQTKANM GSSVSYIMAANHAASVVGAGLAISAERADCEARCARIAREESLLEVPGEENACEKKVAG EKAKTFTRIKYALLTMLEKFLECVADVFKLVPLPITMGIRAIVAAGCTFTSAIIGLCTFCAR A

SEQ ID NO: 29 (groEL-like hsp60 protein)
MVAKNIKYNEEARKKIQKGVKTLAEAVKVTLGPKGRHVVIDKSFGSPQVTKDGVTVAK
EVELADKHENMGAQMVKEVASKTADKAGDGTTTATVLAEAIYTEGLRNVTAGANPMD
LKRGIDKAVKVVVDQIRKISKPVQHHKEIAQVATISANNDAEIGNLIAEAMEKVGKNGSI
TVEEAKGFETVLDIVEGMNFNRGYLSSYFATNPETQECVLEDALVLIYDKKISGIKDFLPV
LQQVAESGRPLLIIAEDIEGEALATLVVNRIRGGFRVCAVKAPGFGDRRKAMLEDIAILTG
GQLISEELGMKLENANLAMLGKAKKVIVSKEDTTIVEGMGEKEALEARCESIKKQIEDSS
SDYDKEKLQERLAKLSGGVAVIRVGAATEIEMKEKKDRVDDAQHATIAAVEEGILPGGG
TALIRCIPTLEAFLPMLTNEDEQIGARIVLKALSAPLKQIAANAGKEGAIIFQQVMSRSANE
GYDALRDAYTDMLEAGILDPAKVTRSALESAASVAGLLLTTEALIAEIPEEKPAAAPAMP
GAGMDY

SEQ ID NO: 30 (omp2/omcB)
MRIGDPMNKLIRRAVTIFAVTSVASLFASGVLETSMAESLSTNVISLADTKAKDNTSHKS
KKARKNHSKETPVDRKEVAPVHESKATGPKQDSCFGRMYTVKVNDDRNVEITQAVPEY

PCT/US2004/020491 WO 2005/002619

ATVGSPYPIEITATGKRDCVDVIITQQLPCEAEFVRSDPATTPTADGKLVWKIDRLGQGEK SKITVWVKPLKEGCCFTAATVCACPEIRSVTKCGQPAICVKQEGPENACLRCPVVYKINI VNQGTATARNVVVENPVPDGYAHSSGQRVLTFTLGDMQPGEHRTITVEFCPLKRGRAT NIATVSYCGGHKNTASVTTVINEPCVQVSIAGADWSYVCKPVEYVISVSNPGDLVLRDV VVEDTLSPGVTVLEAAGAQISCNKVVWTVKELNPGESLQYKVLVRAQTPGQFTNNVVV KSCSDCGTCTSCAEATTYWKGVAATHMCVVDTCDPVCVGENTVYRICVTNRGSAEDTN VSLMLKFSKELQPVSFSGPTKGTITGNTVVFDSLPRLGSKETVEFSVTLKAVSAGDARGE AILSSDTLTVPVSDTENTHIY

SEQ ID NO: 31

MFRYTLSRSLFFILALFFCSACDSRSMITHGLSGRDANEIVVLLVSKGVAAQKVPQAASST GGSGEQLWDISVPAAQITEALAILNQAGLPRMKGTSLLDLFAKQGLVPSEMQEKIRYQEG LSEQMATTIRKMDGIVDASVQISFSPEEEDQRPLTASVYIKHRGVLDNPNSIMVSKIKRLV ASAVPGLCPENVSVVSDRASYSDITINGPWGLSDEMNYVSVWGIILAKHSLTKFRLVFYF LILLLFILSCGLLWVIWKTHTLISALGGTKGFFDPAPYSQLSFTQNKPAPKETPGAAEGAE AQTASEQPSKENAEKQEENNEDA

SEQ.ID NO.32 (CT600)
MRKTIFKAFNLLFSLLFLSSCSYPCRDWECHGCDSARPRKSSFGFVPFYSDEEIQQAFVED FDSKEEQLYKTSAQSTSFRNITFATDSYSIKGEDNLTILASLVRHLHKSPKATLYIEGHTDE RGAAAYNLALGARRANAVKQYLIKQGIAADRLFTISYGKEHPVHPGHNELAWQQNRRT **EFKIHAR** 

**SEQ ID NO 33 (CT541)** 

 ${\bf MKNILSWMLMFAVALPIVGCDNGGGSQTSATEKSMVEDSALTDNQKLSRTFGHLLSRQ}$ LSRTEDFSLDLVEVIKGMQSEIDGQSAPLTDTEYEKQMAEVQKASFEAKCSENLASAEKF LKENKEKAGVIELEPNKLQYRVVKEGTGRVLSGKPTALLHYTGSFIDGKVFDSSEKNKEP ILLPLTKVIPGFSQGMQGMKEGEVRVLYIHPDLAYGTAGQLPPNSLLIFEVKLIEANDDNV SVTE"

**SEQ ID NO 34 (CT623)** MKKYFYKGFVGALLLACGSTNLAFAQASSMDSQLWSVEDLDSYLSSKGFVETRKRDGV LRLAGDVRARWIYAKEDLETTQTPAKPMLPTNRYRSEFNLYVDYTAANSWMTSKMNW VTIAGGESSAAGLDINRAFLGYRFYKNPETQAEVFAEIGRSGLGDIFDSDVQFNSNFDGIH LYAARRISEKLPFTMIVHGGPFVVNMAEKEYAWVVEAILNKLPGNFVVKTSVVDWNTL TAKTNDPADASAAQPAKPNTKYDYLVWQWLVGKSTAMPWFNGQTKNLYTYGAYLFN PLAEIPENWKQSTTPTTKITNGKENHAWFIGCSLGGVRRAGDWSATVRYEYVEALAIPEI DVAGIGRGNQMKYWFAQAIKQGLDPKESNGFTNYKGVSYQFVMGLTDSVSFRAYAAY SKPANDNLGSDFTYRKYDLGLISSF

**SEQ ID NO 35 (CT700)** MWLIVASTLLACLAMALVFKAYRHVISFRSYVNQVIRDVRLSVDLKEWAVAEMRLAPIL KKRQYRRKYLFEYIRILRELERFEEAEKLLGEAKKLKLAGAHFFLEVAHKAFRHGAYKE AAHAFSLLSAELMGEREVARYTISLVYLGEVDAACRIIEPWIGPLAHQEVFISVGHIYFAT KRYADAIDFYRRARSLGSCPIDVLYNLAHSLRICGQYVDAGMLFRELLGDPVYKDEAMF NIGLCEQKLGNSKKALLIYQNSELWVRGDALMMRYAALAAADQQDYQLAEHCWTLAF RCQSYADDWNCCVHYGLALCHLKKYAEAEKVYLRVIQKTPDCLVACKALAWLAGVG HATMISAREGIAYAKRALQIKRSPEVLELLSACEAREGNFDVAYDIQAILAERDTTAKER ERRSQILKNLRQKLPIDQQHIVEVSLLLAA

**SEQ ID NO 36 (CT266)** 

PCT/US2004/020491

MLVESQLGLEDVLEAFSERNFDIQSKSFIESFQDKKLRRTVIQRFLHHPLLHIHDIARAAYL LAALEEGVDLGYQFLCMHQTQSGAALLFRRAGFLWGGLPYPGEHAEMAMLLSRIAEFY DTSYEQVQKMIAFQHALFSHERNIFPALWSQEGSRSNQEKTAVSKLLFCQKEARIEDQFT LTDMSLGFWMRRTPSFSAYVSGSGCKSGVGAFLIGDVGVLNYGPCVGDPGECLGFGLC GQVKEFSCQEKDEEVSISFAGALSQPSSRRTGFSYLQDALFSTNSCYCIDITEQKCHVASS LDRENQDAFFAIFCKGSQCQVCNGPKLRTGSPDSYKGPAYDVLIKGEKETVRILSSSPHM EIFSLQGKDRFWGSNFLINLPYTQNSINILFEKA

**SEQ ID NO 37 (CT077)** 

MGKFFASYLLILAPFFLQSCSAPSRTTLEGVRMTIPYRIVFGEALSPDAFQQAQKEIDRVF DHIDQTFNNWNPLSEISRINRTTKQTPIPLSPALFAFLCEIDHFHAFSDGRFDPTLGALKSL WLLHLKSHTIPSQELQHLYKHSSGWHLISLDKTQQTLRKLSPLVQLDLCGTVKGFAVDL LGTACAQFCQNYYVEWGGEIKTKGKHPSGRSWAVASSATPEILHLHDHAIATSGSQYQR WHVDNKTYTHILDPLTGTPLEDSSHPILAVSVINESCAFADAMATALTTFSSKQEALDWA **NKKHLCAYITDKNVS** 

**SEQ ID NO 38 (CT456)** 

MTNSISGYQPTVTTSTSSTTSASGASGSLGASSVSTTANATVTQTANATNSAATSSIQTTG ETVVNYTNSASAPNVTVSTSSSSTQATATSNKTSQAVAGKITSPDTSESSETSSTSSSDHIP SDYDDVGSNSGDISNNYDDVGSNNGDISSNYDDAAADYEPIRTTENTYESIGGSRTSGPEN TSGGAAAALNSLRGSSYSNYDDAAADYEPIRTTENIYESIGGSRTSGPENTSGGAAAALN SLRGSSYSNYDDAAADYEPIRTTENIYESIGGSRTSGPENTSDGAAAAALNSLRGSSYTTG PRNEGVFGPGPEGLPDMSLPSYDPTNKTSLLTFLSNPHVKSKMLENSGHFVFIDTDRSSFI LVPNGNWDQVCSIKVQNGKTKEDLDIKDLENMCAKFCTGFSKFSGDWDSLVEPMVSAK AGVASGGNLPNTVIINNKFKTCVAYGPWNSQEASSGYTPSAWRRGHRVDFGGIFEKAND FNKINWGTQAGPSSEDDGISFSNETPGAGPAAAPSPTPSSIPIINVNVNVGGTNVNIGDTNV NTTNTTPTTQSTDASTDTSDIDDINTNNQTDDINTTDKDSDGAGGVNGDISETESSSGDDS GSVSSSESDKNASVGNDGPAMKDILSAVRKHLDVVYPGENGGSTEGPLPANQTLGDVIS DVENKGSAQDTKLSGNTGAGDDDPTTTAAVGNGAEEITLSDTDSGIGDDVSDTASSSGD ESGGVSSPSSESNKNTAVGNDGPSGLDILAAVRKHLDKVYPGDNGGSTEGPLQANQTLG DIVQDMETTGTSQETVVSPWKGSTSSTESAGGSGSVQTLLPSPPPTPSTTTLRTGTGATTT SLMMGGPIKADIITTGGGGRIPGGGTLEKLLPRIRAHLDISFDAQGDLVSTEEPQLGSIVNK FRQETGSRGILAFVESAPGKPGSAQVLTGTGGDKGNLFQAAAAVTQALGNVAGKVNLAI QGQKLSSLVNDDGKGSVGRDLFQAAAQTTQVLSALIDTVG

**SEQ ID NO 39 (CT165)** 

MFQPETVPSNRSTETTPQNIEVYNDRNFTNHTTEDVIRIGERLQRQFYNMTEESRVPFTTS PSHHTGNWKTAFLYNLSQVVAHIFPSTVQPIRVKPTRIPPSPTPPPEGTTTAETSTSENKVT TISKEQEVTTKPLLVRERRSLLHSQ

**SEQ ID NO 40 (CT713)** 

MSSKLVNYLRLTFLSFLGIASTSLDAMPAGNPAFPVIPGINIEQKNACSFDLCNSYDVLSA LSGNLKLCFCGDYIFSEEAQVKDVPVVTSVTTAGVGPSPDITSTTKTRNFDLVNCNLNTN CVAVAFSLPDRSLSAIPLFDVSFEVKVGGLKQYYRLPMNAYRDFTSEPLNSESEVTDGMIEVQSNYGFVWDVSLKKVIWKDGVSFVGVGADYRHASCPIDYIIANSQANPEVFIADSDG KLNFKEWSVCVGLTTYVNDYVLPYLAFSIGSVSRQAPDDSFKKLEDRFTNLKFKVRKITS SHRGNICIGATNYVADNFFYNVEGRWGSQRAVNVSGGFQF

**SEQ ID NO 41 (CT082)** 

MSISGSGNVSPATPDFDPSILMGRQAASAHAAKEASGASKATETSAAEQQALISSGTELD YVTDLQQSEGKYKKTLDKTSKSPKTKLKGNFSKVRAGTKGFLTGFGTRASRISARKAEN WO 2005/002619 PCT/US2004/020491

NGEGMSMIPSQMEYVKKKGNRVSPEMQNFYLGASGLWSPTSDVSSITENCLGATALSTT PLLTTMQDPVSIEHLSSGEITALASFNPNVRTASLNEQTINAWTEARLGGEMVSTLLDPNI ETSSLLRRAPTVSNEGMVDVSDMGNQTTSLSMEGLVNTVVDDPASAEEEKKTGELSLEE MAAMAKMMAALLSSGQGMAVFIASSTPSSGLTQFPEPKFSGTIPHHFSKKEDNETIWGL DSQIGSIAFDTRRENNASPLPTTSLHEEASYRFPVGEAPLDVNEIPFAVQHSTVFSKETANT EQALIQNESLGEIPVSAEVVGQDTVSSAYQFPSHLGMAVLASVPLSTEDYKTAVEHRKGP GGPPDPLIYQYRNVAVDPAIIFQSPSPFSVSSRFSVQGKPEAVAVYNDDQEEAAGGNRDS DEGKDQEQDKTRETEDAGGDS

## **SEQ ID NO 42 (CT181)**

MLSKFCKLSLSAILLINTLAPSETFSEEGTSGFLGRMKSWILKDKTILSTTEESQTSAIEKVS DLLSWKRYDYTQESGFAIQFPESPEHSEQVIEVPQSDLAIRYDTYVAETPSDSTVYVVSIW EYPEKIDISRPELNLQEGFAGMLYALPESQVLYLKATALQGHKALEFWIACDDVYFRGM LVSVNHTLYQVFMVYKGRSPEILDKEYSTFIQSFKVTKVRNSKKMDIRKRVSL

## **SEQ ID NO 43 (CT050)**

MNDTKNNISSSFWNPNKVVTKVLLKVSETGIESTPGIVKHNQLITQSENPTDPTDAVTFK
YLKENYTKENDPNPGFLPTTGGTMTGDIDMQGNNVTDIVMYTNGQQNPTDDSAVTIGY
LNEKADEIKSNDQITTAVAGLSNINSQISTLHQLLGIAEDPDTVTNPDLLKTSGGTVYEDI
DMSSNTVSDLGTPTNKDTKSAINVEFVQAKITSPQMAFLKNNDTNLSNITVSEYFNWLQ
DPTQAPTPEPDPDPEPAPEPEPDTSDSSGSGSENPADPAPTNPSDSNAQNNPTPSSNGATAS
IRKLAATTTTVPTDTEIAPAAEDPNLPNTTFSEKSPLWEEFFSFSDSSRSEMVIQKTGILTFS
MQGTWENPSSSQTPSTDPISLELTVTPPTTDTPPESPPSPPEAPAPEATPSPTNNNLTASITK
TFSRKYNLSATPSPTPTTPTEPTTITKTLSLSSGQSCTLQIPVQATRSVLKLKYVNPNNNSS
GGSSGSGGSSQPETTPTGITLQSFSWSLVLTPGEITKATSTPSTPSQP

## **SEQ ID NO 44 (CT157)**

MSVQGSSSLKYSDLFKPPEPTSSTDSSKEPPKESAWKVVSHSRGRRRARSNPSPHTSQNTP SPKDSSLVARTDKAATDIFNSAKHKAIETTKRSDQQSRSLHILHLLAENPEPIVFHSAHQT NHNDPQRMLCDAILQANRIITMRIFNIGSPEIIRALIRAVRRNIPVVVSAWNFPNLSNWDR ESELCVELRGNPQICLHKKTTLIDNQLTIIGTANYTKSSFFKDINLTALIQNPALYSLILSDT RGSVSIGSQTISYYPLPFPQSNTKILPIIQEIQKAQRTIKIAMNIFSHTEIFLALEQARLRGVTI TIVINKKESAHTLDILHRISALLLLKSVTTVDSLHAKICLIDNQTLIFGSPNWTYHGMHKN LEDLLIVTPLTPKQIHSIQEIWAFLLKNSSPV

## **SEQ ID NO 45 (CT128)**

MDRSPLFLIIMGAPGSGKGTQSKLLASQLSLLHISSGDLLRDAVSKDTPLSQEIKSYLDQG KLLPDTLVWKLVHEKLDEFQQDTLLRRLSFLSRSENSAILDGFPRTVTQAKLLHEFLSSYF PNYKVILLDISDEEVLNRLTSRYICPACQGIYNEQQGFSSCPKCSVELIRRSDDTLEVILDRI QTYKQETQPVLDYYTEKQKLITIDANAPTQQVFQSILDSLSASLVYQERDCCNCDCDDED

## SEQ ID NO 46 (CT153)

MTKPSFLYVIQPFSVFNPRLGRFSTDSDTYIEEENRLASFIESLPLEIFDIPSFMETAISNSPYI LSWETTKDGALFTILEPKLSACAATCLVAPSIQMKSDAELLEEIKQALLRSSHDGVKYRIT RESFSPEKKTPKVALVDDDIELIRNVDFLGRAVDIVKLDPINILNTVSEENILDYSFTRETA QLSADGRFGIPPGTKLFPKPSFDVEISTSIFEETTSFTRSFSASVTFSVPDLAATMPLQSPPM VENGQKEICVIQKHLFPSYSPKLVDIVKRYKREAKILINKLAFGMLWRHRAKSQILTEGS VRLDLQGFTESKYNYQIQVGSHTIAAVLIDMDISKIQSKSEQAYAIRKIKSGFQRSLDDYH IYQIERKQTFSFSPKHRSLSSTSHSEDSDLDLSEAAAFSGSLTCEFVKKSTQHAKNTVTCST AAHSLYTLKEDDSSNPSEKRLDSCFRNWIENKLSANSPDSWSAFIQKFGTHYIASATFGGI GFQVLKLSFEQVEDLHSKKISLETAAANSLLKGSVSSSTESGYSSYSSTSSSHTVFLGGTV

LPSVHDERLDFKDWSESVHLEPVPIQVSLQPITNLLVPLHFPNIGAAELSNKRESLQQAIR VYLKEHKVDEQGERTTFTSGIDNPSSWFTLEAAHSPLIVSTPYIASWSTLPYLFPTLRERSS ATPIVFYFCVDNNEHASQKILNQSYCFLGSLPIRQKIFGSEFASFPYLSFYGNAKEAYFDNT YYPTRCGWIVEKLNTTQDQFLRDGDEVRLKHVSSGKYLATTPLKDTHGTLTRTTNCEDA **IFIIKKSSGY** 

**SEQ ID NO 47 (CT262)** 

MKKFLLLSLMSLSSLPTFAANSTGTIGIVNLRRCLEESALGKKESAEFEKMKNQFSNSMG KMEEELSSIYSKLQDDDYMEGLSETAAAELRKKFEDLSAEYNTAQGQYYQILNQSNLKR MQKIMEEVKKASETVRIQEGLSVLLNEDIVLSIDSSADKTDAVIKVLDDSFONN

**SEQ ID NO 48 (CT276)** 

MFKRPAKNFFDEVQTLYEDSGANSTSYSIYPQRTERLENHSNIFEPAKPAETRLLSQEEHS QWTDQQEELATQESSFPEEPETTLGEGVSFKGELTFERLLRIDGTFEGILVSKGKIIVGPQG YVKANIELEEAVIAGVVEGNITVTGRVSLQGRAMVTGDIQAGSLCVDEGVRLCGYVSIQ **GAPSNEQEEIDS** 

SEO ID NO 49 (CT296)

MRAVLHLEHKRYFQNHGHILFEGLAPVSDCKQLEAELKLFLKEVAVVKDRHLQRWREN VHRTLPEVQMIVKRVRLDHLAAELTHRSRVALVRDLWVQKQEEIFFDDCDCSVLLCLSG EKAGWGLFFSGEYPQDVFNWGAGDTAIILRFSSAGFPN

**SEQ ID NO 50 (CT372)** 

MQAAHHHYHRYTDKLHRQNHKKDLISPKPTEQEACNTSSLSKELIPLSEQRGLLSPICDFI SERPCLHGVSVRNLKQALKNSAGTQIALDWSILPQWFNPRVSHAPKLSIRDFGYSAHQTV TEATPPCWQNCFNPSAAVTIYDSSYGKGVFQISYTLVRYWRENAATAGDAMMLAGSIN DYPSRQNIFSQFTFSQNFPNERVSLTIGQYSLYAIDGTLYNNDQQLGFISYALSQNPTATYS SGSLGAYLQVAPTASTSLQIGFQDAYNISGSSIKWSNLTKNRYNFHGFASWAPRCCLGSG QYSVLLYVTRQVPEQMEQTMGWSVNASQHISSKLYVFGRYSGVTGHVFPINRTYSFGM ASANLFNRNPQDLFGIACAFNNVHLSASPNTKRKYETVIEGFATIGCGPYLSFAPDFQLYL YPALRPNKQSARVYSVRANLAI

SEO ID NO 51 (CT412)

MNRVIEIHAHYDQRQLSQSPNTNFLVHHPYLTLIPKFLLGALIVYAPYSFAEMELAISGHK QGKDRDTFTMISSCPEGTNYIINRKLILSDFSLLNKVSSGGAFRNLAGKISFLGKNSSASIH FKHININGFGAGVFSESSIEFTDLRKLVAFGSESTGGIFTAKEDISFKNNHHIAFRNNITKGN GGVIQLQGDMKGSVSFVDQRGAIIFTNNQAVTSSSMKHSGRGGAISGDFAGSRILFLNNQ QITFEGNSAVHGGAIYNKNGLVEFLGNAGPLAFKENTTIANGGAIYTSNFKANQQTSPILF SQNHANKKGGAIYAQYVNLEQNQDTIRFEKNTAKEGGGAITSSQCSITAHNTIIFSDNAA GDLGGGAILLEGKKPSLTLIAHSGNIAFSGNTMLHITKKASLDRHNSILIKEAPYKIQLAA NKNHSIHFFDPVMALSASSSPIQINAPEYETPFFSPKGMIVFSGANLLDDAREDVANRTSIF NQPVHLYNGTLSIENGAHLIVQSFKQTGGRISLSPGSSLALYTMNSFFHGNISSKEPLEING LSFGVDISPSNLQAEIRAGNAPLRLSGSPSIHDPEGLFYENRDTAASPYQMEILLTSDKIVDI SKFTTDSLVTNKQSGFQGAWHFSWQPNTINNTKQKILRASWLPTGEYVLESNRVGRAVP NSLWSTFLLLQTASHNLGDHLCNNRSLIPTSYFGVLIGGTGAEMSTHSSEEESFISRLGAT GTSIIRLTPSLTLSGGGSHMFGDSFVADLPEHITSEGIVQNVGLTHVWGPLTVNSTLCAAL DHNAMVRICSKKDHTYGKWDTFGMRGTLGASYTFLEYDQTMRVFSFANIEATNILQRA FTETGYNPRSFSKTKLLNIAIPIGIGYEFCLGNSSFALLGKGSIGYSRDIKRENPSTLAHLAM NDFAWTTNGCSVPTSAHTLANQLILRYKACSLYITAYTINREGKNLSNSLSCGGYVGF

**SEQ ID NO 52 (CT480)** 

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MIDKIRTILVLSLFLLYWSSDLLEKDVKSIKRELKALHEDVLELVRISHQQKNWVQSTDF SVSPEISVLKDCGDPAFPNLLCEDPYVEKVVPSLLKEGFVPKGILRTAQVGRPDNLSPFNG FVNIVRFYELCVPNLAVEHVGKYEEFAPSLALKIEEHYVEDGSGDKEFHIYLRPNMFWEP IDPTLFPKNITLADSFLRPHPVTAHDVKFYYDVVMNPYVAEMRAVAMRSYFEDMVSVR VENDLKLIVRWRAHTVRNEQGEEEKKVLYSAFANTLALQPLPCFVYQHFANGEKIVPED SDPDTYRKDSVWAQNFSSHWAYNYIVSCGAFRFAGMDDEKITLVRNPNYHNPFAALVE KRYTYMKDSTDSLFQDFKAGKVDIAYFPPNHVDNLASFMQTSAYKEQAARGEAILEKNS SDRSYSYIGWNCLSLFFNNRSVRQAMNMLIDRDRIIEQCLDGRGVSVSGPFSLCSPSYNR DVEGWQYSPEEAARKLEEEGWIDADGDGIREKVIDGVVVPFRFRLCYYVKSVTARTIAE YVATVCKEVGIECCLLGLDMADYSQALEEKNFDAILSGWCLGTPPEDPRALWHSEGALE KGSANAVGFCNEEADRIIEQLSYEYDSNKRQALYHRFHEVIHEESPYAFLYSRQYSLVYK EFVKNIFVPTEHODLIPGAQDETVNLSMLWVDKEEGRCSAIS

## **SEQ ID NO 53 (CT548)**

MLKMFWLNSLVFFSLLLSACGYTVLSPHYVEKKFSLSEGIYVCPIEGDSLGDLVSSLSYEL EKRGLHTRSQGTSSGYVLKVSLFNETDENIGFAYTPQKPDEKPVKHFIVSNEGRLALSAK VQLIKNRTQEILVEKCLRKSVTFDFQPDLGTANAHQLALGQFEMHNEAIKSASRILYSQL AETIYQQYYYDLF

## **SEQ ID NO 54 (CT043)**

MSRQNAEENLKNFAKELKLPDVAFDQNNTCILFVDGEFSLHLTYEEHSDRLYVYAPLLD GLPDNPQRRLALYEKLLEGSMLGGQMAGGGVGVATKEQLILMHCVLDMKYAETNLLK AFAQLFIETVVKWRTVCSDISAGREPTVDTMPQMPQGGGGGIQPPPAGIRA

## **SEQ ID NO 55 (CT635)**

MKNNSAQKIIDSIKQILSIYKIDFEPSFGATLTDDNDLDYQMLIEKTQEKIQELDKRSQEIL QQTGMTREQMEVFANNPDNFSPEEWRALENIRSSCNEYKKETEELIKEVTNDIGHSSHKS PTPKKTKSSSQKKSKKKNWIPL

## **SEQ ID NO 56 (CT859)**

MRKIILCSPRGFCAGVIRAIQTVEVALEKWGRPIYVKHEIVHNRHVVDKLREKGAIFIEDL QEVPRNSRVIFSAHGVPPSVREEAEERGLIAIDATCGLVTKVHSAVKMYAKKGYHIILIGK RKHVEIIGIRGEAPDQITVVENIAEVEALPFSAQDPLFYVTQTTLSMDDAADIVAALKARY PRIFTLPSSSICYATQNRQGALRNILPQVDFVYVIGDTQSSNSNRLREVAERRGVTARLVN HPDEVTEEILQYSGNIGITAGASTPEDVVQACLMKLQELIPDLSIEMDLFVEEDTVFQLPK EL

# **SEQ ID NO 57 (CT671)**

MELNKTSESLFSAKIDHNHPRTEAHEPRDQREVRVFSLEGRSSTRQEKADRMPGRTSSRQ ESSKGSEEGAVHESTAGVSSKEEEESKGDGFFTGGNPTSGMALVETPMAVVSEAMVETS TMTVSQVDLQWVEQLVTSTVESLLVADIDGKQLVEIVLDNSNTVPAAFCGANLTLVQTG EEISVSFSNFVDQAQLTEATQLVQQNPKQLVSLVESLKARQLNLTELVVGNVAVSLPTIE KIETPLHMIAATIRHHDQEGDQEGEGRQDQHQGQHQEKKVEEAHI

## **SEQ ID NO 58 (CT016)**

MKVKINDQFICISPYISARWNQIAFIESCDGGTEGGITLKLHLIDGETVSIPNLGQAIVDEVF QEHLLYLESTAPQKNKEEEKISSLLGAVQQMAKGCEVQVFSQKGLVSMLLGGAGSINVL LQHSPEHKDHPDLPTDLLERIAQMMRSLSIGPTSILAKPEPHCNCLHCQIGRATVEEEDAG VSDEDLTFRSWDISQSGEKMYTVTDPLNPEEQFNVYLGTPIGCTCGQPYCEHVKAVLYT

**SEQ ID NO 59 (CT017)** 

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MLIFALSFGADACLCAADLSKAKVEASVGDRAAFSPFTGEIKGNRVRLRLAPHTDSFIIKE LSKGDCLAVLGESKDYYVVAAPEGVRGYVFRTFVLDNVIEGEKVNVRLEPSTSAPILARL SKGTVVKTLGAAQGKWIEIALPKQCVFYVAKNFVKNVGALDLYNQKEGQKKLALDLLS SAMDFADAELQKKIEDIDLDAIYKKMNLAQSEEFKDVPGLQSLVQKALERVQEAFLAKS LEKSSVKVPEIRHKVLEEIAVVSPAVEETPVVTKTEEQKVTTVPVPAPAVVTEPAQDLSSV KGSLLSHYIRKKGFVKASPVIEGRESFERSLFAVWLSLQPEEIRHQLTMESFYRDEQKKKR VLTGELEVYPHIVKNNPGDYLLKNGEDVVAFVYATSIDLSKWLGKSVVLECVSRPNNHF AFPAYTVLSVKEGA

SEQ ID NO 60 (CT043)
MSRQNAEENLKNFAKELKLPDVAFDQNNTCILFVDGEFSLHLTYEEHSDRLYVYAPLLD
GLPDNPQRRLALYEKLLEGSMLGGQMAGGGVGVATKEQLILMHCVLDMKYAETNLLK
AFAQLFIETVVKWRTVCSDISAGREPTVDTMPQMPQGGGGGIQPPPAGIRA

SEQ ID NO 61 (CT548)
MLKMFWLNSLVFFSLLLSACGYTVLSPHYVEKKFSLSEGIYVCPIEGDSLGDLVSSLSYEL
EKRGLHTRSQGTSSGYVLKVSLFNETDENIGFAYTPQKPDEKPVKHFIVSNEGRLALSAK
VQLIKNRTQEILVEKCLRKSVTFDFQPDLGTANAHQLALGQFEMHNEAIKSASRILYSQL
AETIVQQVYYDLF